IREMENTS FOR PATENT APPLIATIONS CONTAINING NOTICE TO COMPLY WITH RE NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821 -1.825 for the following reason(s): This application clearly fails to comply with the requirements of 37 CFR 1.821 Applicant's attention is directed to these regulations, published at 1114 OG 29, - 1.825. May 15, 1990 and at 55 FR 18230, May 1, 1990. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 CFR 1.821(c). A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 CFR 1.821(e). A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached copy of the marked-up "Raw Sequence Listing." The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CFR 1.825(d). The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e). 7. Other: . Applicant must provide: An initial or substitute computer readable form (CRF) copy of the "Sequence An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d)

For questions regarding compliance with these requirements, please contact:

For Rules Interpretation, call (703) 308-1123

For CRF submission help, call (703) 308-4212

For PatentIn software help, call (703) 557-0400

Please return a copy of this notice with your response.

B. Suprtz

PAGE: 1

RAW SEQUENCE LISTING

PATENT APPLICATION US/08/913,430

DATE: 02/03/1999

TIME: 15:56:07

Input Set: H913430.RAW

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

```
<110> Walker, John
1
2
           Lee, Rogan
3
           Dougherty, Stephen W.
                                                                 Does Not Comply
     <120> Antigen Composition Against Mycoplasma
 4
                                                             Corrected Diskette Needed
     <130> U-011415-0
5
     <140> US/08/913,430
     <141> 1997-12-09
7
8
     <150> PCT/AU96/00149
9
     <151> 1996-03-15
     <150> PN 1789
10
     <160> (15) /8 show see rege )
<170> Patent In Ver. 2.0 - beta
11
12
13
```

## ERRORED SEQUENCES FOLLOW

33

34

35

36

37

38

<221> modified\_base

<221> modified base

<222> (3)

<222> (9)

<220>

			_
	14	<210> 16	
	15	<211> 23	
	16	<212> DNA	
	17	<213> Mycoplasma hyopneumoniae	,,
	18	<220>	-
	19	<221> modified_base	
	20	<222> (3)	
	· 21	<213> Mycoplasma hyopneumoniae <220> <221> modified base <222> (3) <223> modified base (Inosine) used in primer sequence <220> <221> modified base <220> <221> modified base <221> modified base <222> (18) <233> modified base (Inosine) used in primer sequence <223	
	22	<220> LL (220)-[225]	/
	23	<221> modified_base	
	24	<222> (18)	
	25	<223> modified base (Inosine) used in primer sequence	
	26	<400> 16 <sub>0</sub>	
E>	27	actiaacgacg agaagcdica ggc	3
	28	<210> 17	_
	29	<211> 23	
	30	<212> DNA	
	31	<213> Mycoplasma hyopneumoniae	
	32	<220>	

<223> modified base (Inosine) used in primer sequence

PAGE:	2		02/03/1999 L5:56:07	
		Input Set: <b>H913</b> 4	130.RAW	
	39 <	223> modified base (Inosine) used in primer sequence		
	40 <	220>		
	41 <	221> modified base		
	42 . <	222> (15)		
	43 <	223> modified base (Inosine) used in primer sequence		
		400> 17,		
E>	45	thagetting tgatingects etc. same even	23	
	46 <	210> 18		
	47 <	211> 20		
	48 <	212> DNA VANT 1001164 1 1/2		
	49 <	211> 20 212> DNA 213> Mycoplasma hyopneumoniae last seguire in file		
	50 <	220>		
	51 <	221> modified_base		
	52 <	222> (18)		
	53 <	223> modified base (Inosine) used in primer sequence		
	54 <	400> 18		
E>	55	aggtcgatga tcttccallcc	20	

PAGE: 3

## VERIFICATION SUMMARY PATENT APPLICATION US/08/913,430

DATE: 02/03/1999 TIME: 15:56:07

Input Set: H913430.RAW

Line	?	Error/Warning	Original Text
	-		
12	E	# of Seq. 15 Not Equal Actual 18	<160> 15
27	E	Wrong Nucleic Acid Designator	aciaacgacg agaagccica ggc
27	E	Wrong Nucleic Acid Designator	aciaacgacg agaagccica ggc
45	E	Wrong Nucleic Acid Designator	ttiagcttig tgatigcctg ctc
45	E	Wrong Nucleic Acid Designator	ttiagcttig tgatigcctg ctc
45	E	Wrong Nucleic Acid Designator	ttiagcttig tgatigcctg ctc
55	E	Wrong Nucleic Acid Designator	aggtcgatga tcttccaicc